RAW SEQUENCE LISTING
ERROR REPORT

BIOTECHNOLOGY
SYSTEMS
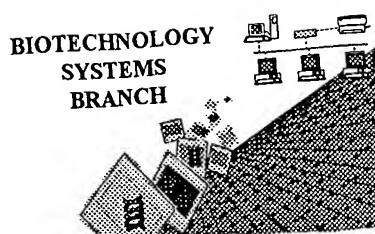**BRANCH**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _10/088,117_

Source: _PCT/10_

Date Processed by STIC: _8/14/2002_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
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2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: __patin21help@uspto.gov__ or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: __patin3help@uspto.gov__ or phone 703-306-4119 (R. Wax)

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Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<__http://www.uspto.gov/ebc/efs/downloads/documents.htm__> , **EFS Submission User Manual** - ePAVE)
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Revised 01/29/2002

PCT10

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/088,117

DATE: 08/14/2002
TIME: 09:31:18

Does Not Comply
Corrected Diskette Needed

Input Set : A:\EP.txt
Output Set: N:\CRF3\08142002\J088117.raw

```
 3 <110> APPLICANT: CNRS
 5 <120> TITLE OF INVENTION: NUCLEOTIDE SEQUENCES DERIVED FROM GENES CODING FOR
 6       TRIMETHYLAMINE N-OXIDE REDUCTASE, AND USES THEREOF,
 7       ESPECIALLY FOR THE DETECTION OF BACTERIA
 9 <130> FILE REFERENCE: WOB 99 AX CNR DORA
11 <140> CURRENT APPLICATION NUMBER: US/10/088,117
12 <141> CURRENT FILING DATE: 2002-07-22
14 <150> PRIOR APPLICATION NUMBER: FR9911543
15 <151> PRIOR FILING DATE: 1999-09-15
17 <160> NUMBER OF SEQ ID NOS: 27
19 <170> SOFTWARE: PatentIn Ver. 2.1
```

ERRORED SEQUENCES

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823 <210> SEQ ID NO: 27
824 <211> LENGTH: 17
825 <212> TYPE: DNA
826 <213> ORGANISM: Artificial sequence
828 <220> FEATURE:
829 <223> OTHER INFORMATION: Description of the artificial sequence:
830       PCR primer
832 <400> SEQUENCE: 27
```

W--> 833 ttngcrtcra artgngc   *see p. 2 for explanation of error*   17
E--> 836 1
E--> 838 1 *delete.*

VARIABLE LOCATION SUMMARY
PATENT APPLICATION: US/10/088,117

DATE: 08/14/2002
TIME: 09:31:20

Input Set : A:\EP.txt
Output Set: N:\CRF3\08142002\J088117.raw

Use of n's or Xaa's(NEW RULES):
Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
in <220> to <223> section, please explain location of n or Xaa, and which
residue n or Xaa represents.

Seq#:22; N Pos. 9
Seq#:24; N Pos. 3,6
Seq#:27; N Pos. 3,15

```
L:11 M:270 C: Current Application Number differs, Replaced Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:768 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:22
L:768 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:22
L:768 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:794 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:24
L:794 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:24
L:794 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
L:833 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:27
L:833 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:27
L:833 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
L:836 M:254 E: No. of Bases conflict, this line has no nucleotides.
M:254 Repeated in SeqNo=27
```